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Team

RAW SEQUENCE LISTING PATENT APPLICATION US/08/794,851 DATE: 08/22/97 TIME: 14:56:26

INPUT SET: S19830.raw

RECEIVED 3 1997 SEP **GROUP 1800**

This Raw Listing contains the General Mages PRED

SEQUENCE LISTING

```
2
 3
     (1)
            General Information:
 4
 5
          (i) APPLICANT: Barany, Francis
                         Barany, George
Hammer, Robert P.
                          Kempe, Maria
 8
 9
                          Blok, Herman
10
                          Zirvi, Monib
11
         (ii) TITLE OF INVENTION: DETECTION OF NUCLEIC ACID SEQUENCE
12
                 DIFFERENCES USING THE LIGASE DETECTION REACTION WITH
13
                 ADDRESSABLE ARRAYS
14
15
        (iii) NUMBER OF SEQUENCES: 16
16
17
18
         (iv) CORRESPONDENCE ADDRESS:
               (A) ADDRESSEE: Nixon, Hargrave, Devans & Doyle LLP
19
               (B) STREET: Clinton Square, P.O. Box 1051
20
21
               (C) CITY: Rochester
               (D) STATE: New York
22
23
               (E) COUNTRY: U.S.A.
24
               (F) ZIP: 14603
25
          (V) COMPUTER READABLE FORM:
26
27
               (A) MEDIUM TYPE: Floppy disk
               (B) COMPUTER: IBM PC compatible
28
               (C) OPERATING SYSTEM: PC-DOS/MS-DOS
29
               (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
30
31
32
         (vi) CURRENT APPLICATION DATA:
33
               (A) APPLICATION NUMBER: US 08/794,851
34
               (B) FILING DATE: 04-FEB-1997
35
               (C) CLASSIFICATION:
36
37
        (vii) PRIOR APPLICATION DATA:
               (A) APPLICATION NUMBER: US 60/011,359
38
39
               (B) FILING DATE: 09-FEB-1996
40
       (viii) ATTORNEY/AGENT INFORMATION:
41
              (A) NAME: Goldman, Michael L.
42
               (B) REGISTRATION NUMBER: 30,727
43
              (C) REFERENCE/DOCKET NUMBER: 19603/461
44
45
         (ix) TELECOMMUNICATION INFORMATION:
46
```

RAW SEQUENCE LISTING PATENT APPLICATION US/08/794,851

DATE: 08/22/97 TIME: 14:56:28

INPUT SET: S19830.raw

| | | 1111 U1 DE1. D17030.7417 |
|----------|--|--------------------------|
| 47 48 | (A) TELEPHONE: (716) 263-1304 (B) TELEFAX: (716) 263-1600 | |
| 49 50 | | |
| 51 52 | (2) INFORMATION FOR SEQ ID NO:1: | |
| 53 | (i) SEQUENCE CHARACTERISTICS: | |
| 54 | (A) LENGTH: 10 base pairs | |
| 55 56 | (B) TYPE: nucleic acid (C) STRANDEDNESS: single | |
| 57 | (D) TOPOLOGY: linear | |
| 58 | (5) 10.02021 22.002 | |
| 59 | (ii) MOLECULE TYPE: cDNA | |
| 60 | | |
| 61 62 | | |
| 63 | | |
| 64 | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1: | |
| 65 | | |
| 66 | CACACACA | 10 |
| 67 68 | (2) INFORMATION FOR SEQ ID NO:2: | |
| 69 | (2) INFORMATION FOR BEQ ID NO.2. | |
| 70 | (i) SEQUENCE CHARACTERISTICS: | |
| 71 | (A) LENGTH: 24 base pairs | |
| 72 | (B) TYPE: nucleic acid | |
| 73 74 | <pre>(C) STRANDEDNESS: single (D) TOPOLOGY: linear</pre> | |
| 75 | (b) forologi. Timedi | |
| 76 | (ii) MOLECULE TYPE: cDNA | |
| 77 | | |
| 78 79 | | |
| 80 | | |
| 81 | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2: | |
| 82 | | |
| 83 84 | TGCGGGTACA GCACCTACCT TGCG | 24 |
| 85 | (2) INFORMATION FOR SEQ ID NO:3: | |
| 86 | (2) Intolumination for DIV ID Notes | |
| 87 | (i) SEQUENCE CHARACTERISTICS: | |
| 88 | (A) LENGTH: 24 base pairs | |
| 89 90 | (B) TYPE: nucleic acid (C) STRANDEDNESS: single | |
| 91 | (D) TOPOLOGY: linear | |
| 92 | . , | |
| 93 | (ii) MOLECULE TYPE: cDNA | |
| 94 95 | * | |
| 95 96 | 4 | |
| 97 | f i | |
| 98 | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3: | |
| 99 | | |

RAW SEQUENCE LISTING PATENT APPLICATION US/08/794,851

DATE: 08/22/97 TIME: 14:56:31

INPUT SET: S19830.raw ATCGGGTAGG TAACCTTGCG TGCG (2) INFORMATION FOR SEQ ID NO:4: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 24 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: cDNA (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4: CAGCGGTAGA CCACCTATCG TGCG (2) INFORMATION FOR SEQ ID NO:5: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 24 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: cDNA (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5: GGTAGGTAAC CTACCTCAGC TGCG (2) INFORMATION FOR SEQ ID NO:6: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 24 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: cDNA (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6: GACCGGTATG CGACCTGGTA TGCG

RAW SEQUENCE LISTING PATENT APPLICATION US/08/794,851

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DATE: 08/22/97 TIME: 14:56:33

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| 153 154 | (2) | INFORMATION FOR SEQ ID NO:7: | * |
|------------|-------|---|----------|
| 155 | | (i) SEQUENCE CHARACTERISTICS: | |
| 156 | | (A) LENGTH: 24 base pairs | |
| 157 | | (B) TYPE: nucleic acid | |
| 158 | | (C) STRANDEDNESS: single | |
| 159 | | (D) TOPOLOGY: linear | |
| 160 | | (b) Toronour. Integr | |
| 161 | | (ii) MOLECULE TYPE: cDNA | |
| 162 | | (11) MODECODE 11FB. CDIR | |
| 163 | | | |
| 164 | | | |
| 165 | | | |
| 166 | | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7: | |
| 167 | | (XI) SEQUENCE DESCRIPTION: SEQ ID NO:/: | |
| | a mar | GGGTAGG TAACCTTGCG TGCG | 24 |
| 169 | AICC | GGGTAGG TAACCTTGCG TGCG | 24 |
| | | THEODMARION ROD CEO ID NO. 0. | |
| | (2) | INFORMATION FOR SEQ ID NO:8: | |
| 171 | | (1) CHOURNOR CUARACTERISTICS. | |
| 172 | | (i) SEQUENCE CHARACTERISTICS: | |
| 173 | | (A) LENGTH: 24 base pairs | |
| 174 | | (B) TYPE: nucleic acid | |
| 175 | | (C) STRANDEDNESS: single | |
| 176 | | (D) TOPOLOGY: linear | |
| 177 | | | |
| 178 | | (ii) MOLECULE TYPE: cDNA | |
| 179 | | | |
| 180 | | | |
| 181 | | | |
| 182 | | | |
| 183 | | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8: | |
| 184 | | | |
| | GGT | AGGTAAC CTACCTCAGC TGCG | 24 |
| 186 | | | |
| | (2) | INFORMATION FOR SEQ ID NO:9: | |
| 188 | | | |
| 189 | | (i) SEQUENCE CHARACTERISTICS: | |
| 190 | | (A) LENGTH: 24 base pairs | * |
| 191 | | (B) TYPE: nucleic acid | |
| 192 | | (C) STRANDEDNESS: single | |
| 193 | | (D) TOPOLOGY: linear | |
| 194 | | | |
| 195 | | (ii) MOLECULE TYPE: cDNA | |
| 196 | | | |
| 197 | | | |
| 198 | | | |
| 199 | | | • |
| 200 | | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9: | |
| 201 | | | Š |
| | ATC | GGGTAGG TAACCTTGCG TGCG | 24 |
| 203 | _ | | B |
| | (2) | INFORMATION FOR SEQ ID NO:10: | • |
| 205 | , , | · · · · · · · · · · · · · · · · · · · | |

RAW SEQUENCE LISTING PATENT APPLICATION US/08/794,851

DATE: 08/22/97 TIME: 14:56:35

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| | | 11(1 01 02)1. 02/00/1140 |
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| 206 207 208 | (i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 24 base pairs(B) TYPE: nucleic acid | |
| 209 210 | (C) STRANDEDNESS: single(D) TOPOLOGY: linear | |
| 211 212 | (ii) MOLECULE TYPE: cDNA | |
| 213 214 | | |
| 215 216 | | |
| 217 | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: | |
| 218 | , , - | |
| 219 | GGTAGGTAAC CTACCTCAGC TGCG | 24 |
| 220 | | |
| 221 | (2) INFORMATION FOR SEQ ID NO:11: | |
| 222 223 | (i) SEQUENCE CHARACTERISTICS: | |
| 224 | (A) LENGTH: 24 base pairs | |
| 225 | (B) TYPE: nucleic acid | |
| 226 | (C) STRANDEDNESS: single | |
| 227 | (D) TOPOLOGY: linear | |
| 228 | | |
| 229 230 | (ii) MOLECULE TYPE: cDNA | |
| 230 | | • |
| 232 | | |
| 233 | | |
| 234 | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: | |
| 235 | | |
| 236 | ATCGGGTAGG TAACCTTGCG TGCG | 24 |
| 237 238 | (2) INFORMATION FOR SEQ ID NO:12: | |
| 239 | (2) INFORMATION FOR SEQ ID NO.12. | |
| 240 | (i) SEQUENCE CHARACTERISTICS: | |
| 241 | (A) LENGTH: 24 base pairs | |
| 242 | (B) TYPE: nucleic acid | |
| 243 | (C) STRANDEDNESS: single | |
| 244 245 | (D) TOPOLOGY: linear | |
| 246 | (ii) MOLECULE TYPE: cDNA | |
| 247 | (11) 110120012 111121 001111 | |
| 248 | | |
| 249 | | |
| 250 | (!) GEOVERNOE REGORDERAN, AND TO WO IS | |
| 251 252 | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:12: | . 6 |
| 252 253 | CAGCACCTGA CCATCGATCG CAGC | 24 |
| 254 | The state of the s | 4 |
| 255 | (2) INFORMATION FOR SEQ ID NO:13: | • |
| 256 | | • |
| 257 | (i) SEQUENCE CHARACTERISTICS: | 7 |
| 258 | (A) LENGTH: 24 base pairs | |

SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/08/794,851

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Original Text